## SEQUENCE LISTING

<110> Fred Hutchinson Cancer Research Center Stoddard, Barry L.
Pratt, Kathleen
Fujikawa, Kazuo
Davie, Earl W.

<120> CRYSTAL OF A TRUNCATED PROTEIN CONSTRUCT CONTAINING A COAGULATION FACTOR VIII C2 DOMAIN IN THE PRESENCE OR ABSENCE OF A BOUND LIGAND AND METHODS OF USE THEREOF

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<150> 60/148,907

<151> 1999-08-13

<160> 12

<170> PatentIn Ver. 2.1

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<213> Homo sapiens

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Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln Ala Glu Val 65 70 75 80

Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser His Pro Val 85 90 95

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Pro Gln Val Asn Asn Xaa Lys Glu Trp Leu Gln Val Asp Phe Gln Lys
Thr Xaa Lys Val Thr Gly Val Thr Xaa Xaa Gly Val Lys Ser Leu Leu
Thr Ser Met Tyr Val Lys Glu Phe Leu Xaa Ser Ser Ser Gln Asp Gly
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His Gln Trp Thr Leu Phe Phe Gln Asn Gly Lys Val Lys Val Xaa Gln
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Pro Gln Val Asn Asp Pro Lys Gln Trp Leu Gln Val Asp Leu Gln Lys
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Thr Met Lys Val Thr Gly Ile Ile Thr Gln Gly Val Lys Ser Leu Phe 65 70 75 80

Thr Ser Met Phe Val Lys Glu Phe Leu Ile Ser Ser Ser Gln Asp Gly
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His His Trp Thr Gln Ile Leu Tyr Asn Gly Lys Val Lys Val Phe Gln
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35 40 45

Pro Gln Ala Asn Asn Pro Lys Glu Trp Leu Gln Val Asp Phe Arg Lys 50 55 60

Thr Met Lys Val Thr Gly Ile Thr Thr Gln Gly Val Lys Ser Leu Leu 65 70 75 80

Ile Ser Met Tyr Val Lys Glu Phe Leu Ile Ser Ser Gln Asp Gly
85 90 95

His Asn Trp Thr Leu Phe Leu Gln Asn Gly Lys Val Lys Val Phe Gln
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Gly Asn Arg Asp Ser Ser Thr Pro Val Arg Asn Arg Leu Glu Pro Pro 115 120 125

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Pro Arg Val Ser Ser Ala Glu Glu Trp Leu Gln Val Asp Leu Gln Lys 50 55 60

Thr Val Lys Val Thr Gly Ile Thr Thr Gln Gly Val Lys Ser Leu Leu 65 70 75 80

Ser Ser Met Tyr Val Lys Glu Phe Leu Val Ser Ser Ser Gln Asp Gly 85 90 95

Arg Arg Trp Thr Leu Phe Leu Gln Asp Gly His Thr Lys Val Phe Gln
100 105 110

Gly Asn Gln Asp Ser Ser Thr Pro Val Val Asn Ala Leu Asp Pro Pro 115 120 125

Leu Phe Thr Arg Tyr Leu Arg Ile His Pro Thr Ser Trp Ala Gln His 130 135 140

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Gln Ala Lys Ala Asn Asn Asn Lys Gln Trp Leu Glu Ile Asp Leu Leu 50 55 60

Lys Ile Lys Lys Ile Thr Ala Ile Ile Thr Gln Gly Cys Lys Ser Leu 65 70 75 80

Ser Ser Glu Met Tyr Val Lys Ser Tyr Thr Ile His Tyr Ser Glu Gln 85 90 95

Gly Val Glu Trp Lys Pro Tyr Arg Leu Lys Ser Ser Met Val Asp Lys
100 105 110

Ile Phe Glu Gly Asn Thr Asn Thr Lys Gly His Val Lys Asn Phe Phe 115 120 125

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Arg Leu His Tyr Ser Gly Ser Ile Asn Ala Trp Ser Thr Lys Glu Trp
35 40 45

Ile Lys Val Asp Leu Leu Ala Pro Met Ile Ile His Gly Ile Lys Thr
50 55 60

Gln Gly Ala Arg Gln Lys Phe Ser Ser Leu Tyr Ile Ser Gln Phe Ile 65 70 75 80

Ile Met Tyr Ser Leu Asp Gly His His Trp Gln Thr Tyr Arg Gly Asn

85 90 95

Ser Thr Gly Thr Leu Met Val Phe Gln Gly Asn Val Asp Ser Ser Gly 100 105 110

Ile Lys His Asn Ile Phe Asn Pro Pro Ile Ile Ala Arg Tyr Ile Arg

Leu His Pro Thr His Tyr Ser Ile Arg Ser Thr Leu Arg Met Glu Leu 130 135 140